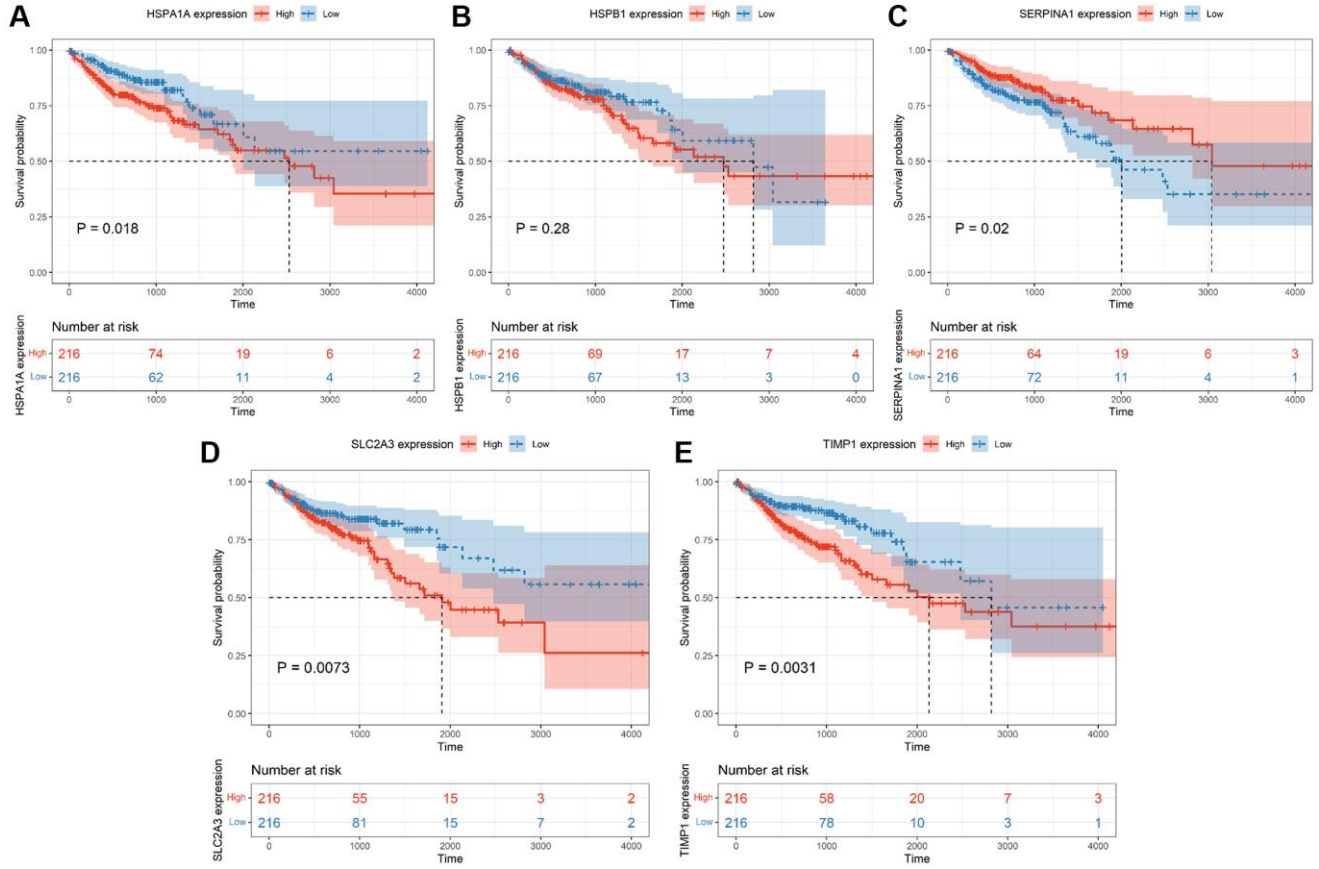
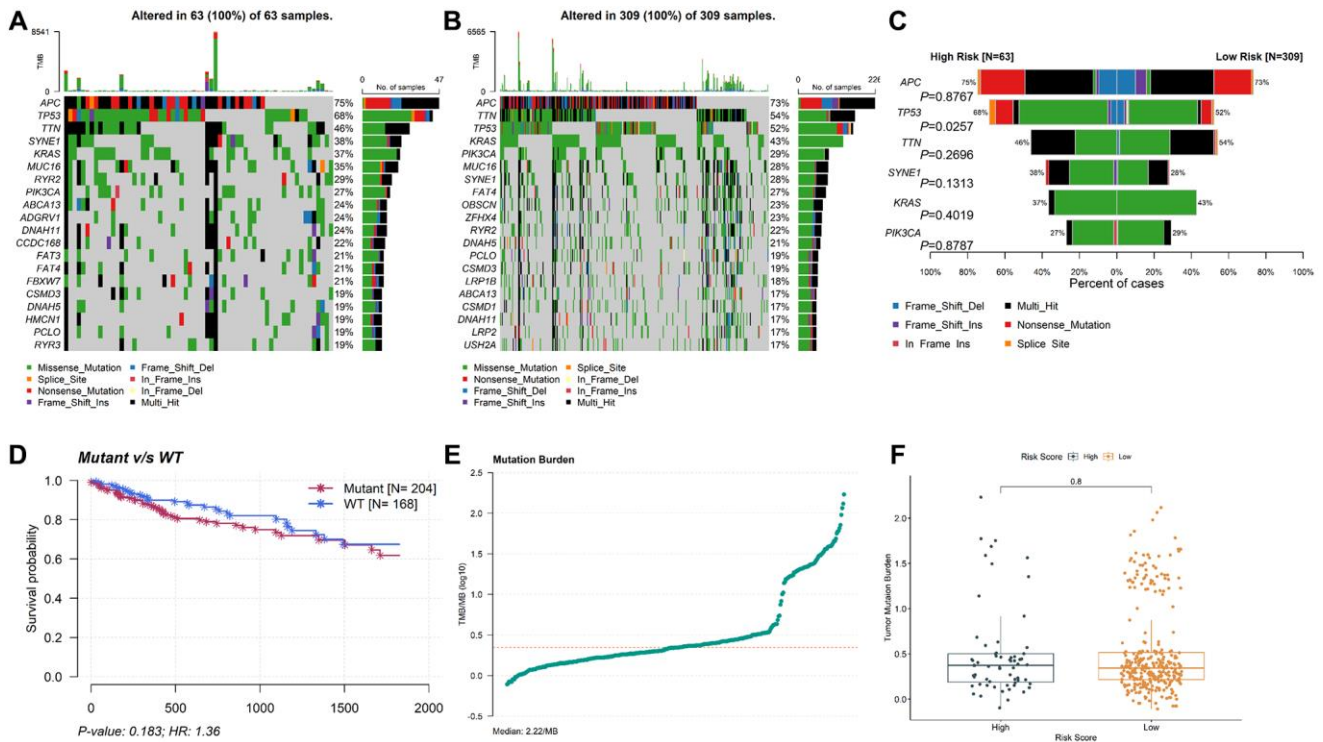


SUPPLEMENTARY FIGURES



Supplementary Figure 1. Kaplan–Meier analysis of each TALRG mRNA expression, including HSPA1A, HSPB1, SERPINA1, SLC2A3, and TIMP1 (A–E).



Supplementary Figure 2. Differences in mutation profiles between high- and low-risk groups. Waterfall plot of the top 20 mutated genes in high- (A) and low-risk groups (B). (C) Comparison of the mutation rate between the high- and low-risk groups. (D) Kaplan-Meier analysis of TP53 mutation status. (E) The TMB profiles of CRC patients. (F) Comparison of TMB values between the high- and low-risk groups.